

FIGURE 1

1 GAATTCAAGA CCAGCCTGGA CAACTTGGAA GAACCCGGTC TCTACAAAAA ATACAAAATT  
 61 AGCTGGGATT GGGTGCCTG GCTCATGCCT ATAATCCCAG CACTTTGGGA GCCTGAGGTG  
 121 GGTGGATCAC CTGAAGTCAG GAGTTCAGA CTAGCCTGGC CAACATGGTG AAACCCCTATC  
 181 TCTACTGAAA ATACAAAAAG CTAGACGTGG TGGCACACAC CTGTAATCCC AGCTACTTAG  
 241 GAGGCTGAGG CAGGAGAATT GCTTGAAGCC TAGAGGTGAA GGTGTAGTG AGCCGAGATT  
 301 GCATCATTCG ACAATGGAGG GGAGCCACCA GCCTGGGCAA CAAGAGGAAA TCTCCGTCTC  
 361 CAAAAA AAAA AAAA AAAGAATTAG GCTGGGTGGT GCCTGTAGTC CCAGCTACTT  
 421 GGGAGGCAGG GGGTCCACTT GATGTCGAGA CTGCAGTGAG CCATGATCCT GCCACTGCAC  
 481 TCCGGCCTGG GCAACAGAGT GAGACCCTGT CTAAAGAAAA AAAAAATAAA GCAACATATC  
 541 CTGAACAAAG GATCCTCCAT AACGTTCCCA CCAGATTCTT AATCAGAAAC ATGGAGGCCA  
 601 GAAAGCAGTG GAGGAGGACG ACCCTCAGGC AGCCCGGGAG GATGTTGTCA CAGGCTGGGG  
 661 CAAGGGCCTT CCGCTACCA ACTGGGAGCT CTGGGAACAG CCCTGTTGCA AACAAGAAGC  
 721 CATAGCCCGG CCAGAGCCCA GGAATGTGGG CTGGGCTGGG AGCAGCCTCT GGACAGGAGT  
 781 GGTCCCATCC AGGAAACCTC CGGCATGGCT GGGAAAGTGG GTACTTGGTG CCGGGTCTGT  
 841 ATGTGTGTGT GACTGGTGTG TGTGAGAGAG AATGTGTGCC CTAAGTGTCA GTGTGAGTCT  
 901 GTGTATGTGT GAATATTGTC TTTGTGTGGG TGATTTTCTG CGTGTGTAAT ACCATCAGGT  
 961 CAAGTGTGAA CAAGTGGACA AGTGTCTGGG AGTGGACAAG AGATCTGTGC ACCATCAGGT  
 1021 GTGTGCATAG CGTCTGTGCA TGTCAAGAGT GCAAGGTGAA GTGAAGGGAC CAGGCCCATG  
 1081 ATGCCACTCA TCATCAGGAG CTCTAAGGCC CCAGGTAAGT GCCAGTGACA GATAAGGGTG  
 1141 CTGAAGGTCA CTCTGGAGTG GGCAGGTGGG GGTAGGGAAA GGGCAAGGCC ATGTTCTGGA  
 1201 GGAGGGGTTG TGACTACATT AGGGTGTATG AGCCTAGCTG GGAGGTGGAT GGCCTGGTCC  
 1261 ACTGAAACCC TGGTTATCCC AGAAGGCTTT GCAGGCTTCA GGAGCTTGGG GTGGGGAGAG  
 1321 GGGGTGACTT CTCCGACCAG GCCCTCCAC CGGCCTACCC TGGGTAAGGG CCTGGAGCAG  
 1381 GAAGCAGGGG CAAGAACCTC TGGAGCAGCC CATAACCGCC CTGGCCTGAC TCTGCCACTG  
 1441 GCAGCACAGT CAACACAGCA GGTTCACCTA CAGCAGAGGG CAAAGGCCAT CATCAGCTCC  
 1501 CTTTATAAGG GAAGGGTCAC GCGCTCGGTG TGCTGAGAGT GTCTGCCTG GTCCTCTGTG  
 1561 CCTGGTGGGG TGGGGGTGCC AGGTGTGTCC AGAGGAGCCC ATTTGGTAGT GAGGCAGGTA  
 1621 TGGGGCTAGA AGCACTGGTG CCCCTGGCCG TGATAGTGGC CATCTTCCTG CTCCTGGTGG  
 1681 ACCTGATGCA CCGGCGCCAA CGCTGGGCTG CACGCTACCC ACCAGGCCCC CTGCCACTGC  
 1741 CCGGGCTGGG CAACCTGTCT CATGTGGACT TCCAGAACAC ACCATACTGC TTCGACCAGG  
 1801 TGAGGGAGGA GGTCTTGGAG GGCGGCAGAG GTGCTGAGGC TCCCCCTACCA GAAGCAAAAC  
 1861 TGGATGGTGG GTGAAACCAC AGGCTGGACC AGAAGCCAGG CTGAGAAGGG GAAGCAGGTT  
 1921 TGGGGGACGT CCTGGAGAAG GGCATTTATA CATGGCATGA AGGACTGGAT TTTCCAAAGG  
 1981 CCAAGGAAGA TAGGGCAAG GGCCTGGAGG TGGAGCTGGA CTTGGCAGTG GGCATGCAAG  
 2041 CCCATTGGGC AACATATGTT ATGGAGTACA AAGTCCCCTC TGCTGACACC AGAAGGAAAG  
 2101 GCCTTGGGAA TGAAGATGA GTTAGTCTTG AGTGCCGTTT AAATCACGAA ATCGAGGATG  
 2161 AAGGGGGTGC AGTGACCCGG TTCAAACCTT TTGCACTGTG GGTCTCGGG CCTCACTGCC  
 2221 TCACCGGCAT GGACCATCAT CTGGGAATGG GATGCTAACT GGGGCTCTC GGCAATTTTG  
 2281 GTGACTCTTG CAAGGTCTA CTTGGGTGAC GCATCCAAAC TGAGTTCCTC CATCACAGAA  
 2341 GGTGTGACCC CCACCCCGC CCCACGATCA GGAGGCTGGG TCTCCTCCTT CCACCTGCTC  
 2401 ACTCCTGGTA GCGCCGGGGG TCGTCCAAGG TTCAAATAGG ACTAGGACCT GTAGTCTGGG  
 2461 GTGATCCTGG CTTGACAAGA GGCCTGACC CTCCCTCTGC AGTTGCGGGC CCGCTTCGGG  
 2521 GACGTGTTCA GCCTGCAGCT GGCCTGGACG CCGGTGGTCG TGCTCAATGG GCTGGCGGCC  
 2581 GTGCGCGAGG CGCTGGTGAC CCACGGCGAG GACACCGCCG ACCGCGGCCG TGTGCCCATC  
 2641 ACCCAGATCC TGGGTTTTCG GCCGCGTTCC CAAGGCAAGC AGCGGTGGGG ACAGAGACAG  
 2701 ATTTCCGTGG GACCCGGGTG GGTGATGACC GTAGTCCGAG CTGGGCAGAG AGGGCGCGGG  
 2761 GTCGTGGACA TGAACAGGC CAGCGAGTGG GGACAGCGGG CCAAGAAACC ACCTGCACTA  
 2821 GGGAGGTGTG AGCATGGGGA CGAGGGCGGG GCTTGTGAC AGTGGGCGGG GCCACTGCCG  
 2881 AGACCTGGCA GGAGCCCAAT GGGTGAGCGT GGCGCATTTT CCAGCTGGAA TCCGTTGTCT  
 2941 AAGTGGGGGC GGGGACCGCA CCTGTGCTGT AAGCTCAGTG TGGGTGGCGC GGGGCGCGC  
 3001 GGGTCTTCCC TGAGTGCAAA GGCGGTCAGG GTGGGCAGAG ACGAGGTGGG GCAAAGCCTG  
 3061 CCCAGCCAA GGGAGCAAGG TGGATGCACA AAGAGTGGGC CCTGTGACCA GCTGGACAGA  
 3121 GCCAGGGACT GCGGGAGACC AGGGGGAGCA TAGGGTTGGA GTGGGTGGTG GATGGTGGGG  
 3181 CTAATGCCTT CATGGCCACG CGCACGTGCC CGTCCCACCC CCAGGGGTGT TCCTGGCGCG  
 3241 CTATGGGCCC GCGTGGCGCG AGCAGAGGCG CTTCTCCGTG TCCACCTTGC GCAACTTGGG  
 3301 CCTGGGCAAG AAGTCGCTGG AGCAGTGGGT GACCGAGGAG GCCGCTGCC TTTGTGCCGC  
 3361 CTTGCGCAAC CACTCCGGTG GGTGATGGGC AGAAGGGCAC AAAGCGGGAA CTGGGAAGGC  
 3421 GGGGGACGGG GAAGGCGACC CTTACCCACC ATCTCCACCC CCCAGGACGC CCCTTTCGCC  
 3481 CCAACGGTCT CTTGGACAAA GCCGTGAGCA ACGTGATCGC CTCCCTCACC TCGGGCGGCC  
 3541 GCTTCGAGTA CGACGACCTT CGCTTCCTCA GGCTGCTGGA CCTAGCTCAG GAGGGACTGA  
 3601 AGGAGGAGTC GGGCTTTCTG CGCGAGGTGC GGAGCGAGAG ACCGAGGAGT CTCTGCAGGG  
 3661 CGAGCTCCCG AGAGGTGCCG GGGCTGACT GGGGCCCTCG AAGAGCAGGA TTTGCATAGA  
 3721 TGGGTTTGGG AAAGGACATT CCAGGAGACC CCACTGTAAG AAGGGCTGG AGGAGAGGGG  
 3781 GACATCTCAG ACATGGTCTG GGGAGAGGTG TGCCCGGGTC AGGGGGCACC AGGAGAGGCC  
 3841 AAGGACTCTG TACCTCCTAT CCACGTGAGA GATTTTCGATT TTAGGTTTCT CCTCTGGGCA  
 3901 AGGAGAGAGG GTGGAGGCTG GCACTTGGGG AGGGACTTGG TGAGGTCAAG GGTAAAGACA

3961	GGCAGGCCCT	GGGTCTACCT	GGAGATGGCT	GGGGCCTGAG	ACTTGTCCAG	GTGAACGCAG
4021	AGCACAGGAG	GGATTGAGAC	CCCCTTCTGT	CTGGTGTAGG	TGCTGAATGC	TGTCCTCCGTC
4081	CTCCTGCATA	TCCAGCGCT	GGCTGGCAAG	GTCTACGCT	TCCAAAAGGC	TTTCTTGACC
4141	CAGCTGGATG	AGCTGCTAAC	TGAGCACAGG	ATGACCTGGG	ACCCAGCCCA	GCCCCCCCCA
4201	GACCTGACTG	AGGCCCTTCCT	GGCAGAGATG	GAGAAGGTGA	GAGTGGCTGC	CACGGTGGGG
4261	GGCAAGGGTG	GTGGGTGAG	CGTCCCAGGA	GGAATGAGGG	GAGGCTGGGC	AAAAGGTTGG
4321	ACCAAGTGCAT	CACCCGGCGA	GCCGCATCTG	GGCTGACAGG	TGCAGAATTG	GAGGTCATTT
4381	GGGGGCTACC	CCGTTCTGTC	CCGAGTATGC	TCTCGGCCCT	GCTCAGGCCA	AGGGGAACCC
4441	TGAGAGCAGC	TTCAATGATG	AGAACCTGCG	CATAGTGGTG	GCTGACCTGT	TCTCTGCCCG
4501	GATGGTGACC	ACCTCGACCA	CGCTGGCCTG	GGGCCTCCTG	CTCATGATCC	TACATCCGGA
4561	TGTGCAGCGT	GAGCCCATCT	GGGAAACAGT	GCAGGGGCCG	AGGGAGGAAG	GGTACAGGCG
4621	GGGGCCCATG	AACCTTGCTG	GGACACCCGG	GGCTCCAAGC	ACAGGCTTGA	CCAGGATCCT
4681	GTAAGCCTGA	CCTCCTCCAA	CATAGGAGGC	AAGAAGGAGT	GTCAGGGCCG	GACCCCTTGG
4741	GTGCTGACCC	ATTGTGGGGA	CGCATGTCTG	TCCAGGCCGT	GTCCAACAGG	AGATCGACGA
4801	CGTGATAGGG	CAGGTGCGGC	GACCAGAGAT	GGGTGACCAG	GCTCACATGC	CCTACACCAC
4861	TGCCGTGATT	CATGAGGTGC	AGCGCTTTGG	GGACATCGTC	CCCCTGGGTG	TGACCCATAT
4921	GACATCCCGT	GACATCGAAG	TACAGGGCTT	CCGCATCCCT	AAGGTAGGCC	TGGCGCCCTC
4981	CTCACCCAG	CTCAGCACCA	GCACCTGGTG	ATAGCCCCAG	CATGGCTACT	GCCAGGTGGG
5041	CCCACTCTAG	GAACCTTGGC	CACCTAGTCC	TCAATGCCAC	CACACTGACT	GTCCCCACTT
5101	GGGTGGGGGG	TCCAGAGTAT	AGGCAGGGCT	GGCCTGTCCA	TCCAGAGCCC	CCGTCTAGTG
5161	GGGAGACAAA	CCAGGACCTG	CCAGAATGTT	GGAGGACCCA	ACGCCTGCAG	GGAGAGGGGG
5221	CAGTGTGGGT	GCCTCTGAGA	GGTGTGACTG	CGCCCTGCTG	TGGGGTCCGA	GAGGGTACTG
5281	TGGAGCTTCT	CGGGCGCAGG	ACTAGTTGAC	AGAGTCCAGC	TGTGTGCCAG	GCAGTGTGTG
5341	TCCCCCGTGT	GTTTGGTGGC	AGGGGTCCCA	GCATCCTAGA	GTCCAGTCCC	CACTCTCACC
5401	CTGCATCTCC	TGCCCAGGGA	ACGACACTCA	TCACCAACCT	GTCATCGGTG	CTGAAGGATG
5461	AGGCCGTCTG	GGAGAAGCCC	TTCCGCTTCC	ACCCCGAACA	CTTCTGGGAT	GCCCAGGGCC
5521	ACTTTGTGAA	GCCGGAGGCC	TTCTCTGCCT	TCTCAGCAGG	TGCCCTGTGG	GAGCCCGGCT
5581	CCCTGTCCCC	TTCCGTGGAG	TCTTGCAGGG	GTATCACCCA	GGAGCCAGGC	TCACTGACGC
5641	CCCTCCCCCT	CCCACAGGCC	GCCGTGCATG	CCTCGGGGAG	CCCCTGGCCC	GCATGGAGCT
5701	CTTCTCTCTC	TTTCACTCCC	TGCTGCAGCA	CTTCAGCTTC	TCGGTGCCCA	CTGGACAGCC
5761	CCGGCCAGC	CACCATGGTG	TCTTTGCTTT	CCTGGTGAGC	CCATCCCCCT	ATGAGCTTTG
5821	TGCTGTGCCC	CGCTAGAAATG	GGGTACCTTT	TCCCCAGCCT	GCTCCCTAGC	CAGAGGCTCT
5881	AATGTACAAT	AAAGCAATGT	GGTAGTTCCA	ACTCGGGTCC	CCTCGTCCAG	CCCTCGTTGG
5941	GATCATCTCT	CTCAGGGCAA	CCCCACCCCT	GCCTCATTCC	TGCTTACCCC	ACCGCCTGGC
6001	CGCATTTGAG	ACAGGGGTAC	GTTGAGGCTG	AGCAGATGTC	AGTTACCCCT	GCCCATAATC
6061	CCATGTCCCC	CACGTAGCCCA	ACTCTGACTG	CCCAGATTGG	TGACAAGGAC	TACATTGTCC
6121	TGGCATGTGG	GGAAGGGGCC	AGAATGGGCT	GACTAGAGGT	GTCAAGTCAGC	CTTGGATGTG
6181	GTGGAGAGGG	CAGGACTCAG	CCTGGAGGCC	CATATTTTCC	GCCTAACTCA	GCCCCACCCA
6241	CATCAGGGAC	AGCAGTCCCTG	CCAGCACCAT	CACAACAGTC	ACCTCCCTTC	ATATATGACA
6301	CCCCAAAACG	GAAGACAAAT	CATGGCGTCA	GGGAGCTATA	TGCCAGGGCT	ACCTACCTCC
6361	CAGGGCTCAG	TCCGGCAGGTG	CCAGAACGTT	CCCTGGGAAG	GCCCCATGGA	AGCCCAGGAC
6421	TGAGCCACCA	CCCTCAGCCT	CGTCACCTTA	CCACAGGACT	GGCTACCTCT	CTGGGCCCTC
6481	AGGGATGCTG	CTGTACAGAC	CCCTGACCAG	TGACGAGTTC	GCACTCAGGG	CCAGGCTGGC
6541	GCTGGAGGAG	GACACTTGT	TGGCTCCAAC	CCTAGGTACC	ATCCTCCCAG	TAGGGATCAG
6601	GCAGGGCCCA	CAGGCCCTGCC	CTAGGGAGAG	GAGTCAACCT	TGGACCCATA	AGGCACTGGG
6661	GCGGGCAGAG	AAGGAGGAGG	TGGCATGGGC	AGCTGAGAGC	CAGAGACCTT	GACCCTAGTC
6721	CTTGCTCTGC	CATTACCCCG	TGTGACCCCG	GGCCCCACCT	TCCCCACCCCT	TCCCCACCCCT
6781	GGGCTTCTGT	TTCTTCTCTG	CAACGAGAAG	GCTGCTTAC	CTGCCCCGAG	TCCTGTCTTC
6841	CTGCTCTGCC	TTCTGGGGCT	GTGGCCCTTG	CTGGCCTGGA	GCCCCAACCA	AGGGCAGGGA
6901	CTGCTGTCTT	CCACGTCTGT	CCTCACCGAC	ATAATGGGCT	GGGCTGGGCA	CACAGGCAGT
6961	GCCCCAAGAGT	TTCTAATGAG	CATATGATTA	CCTGAGTCCCT	GGGCAGACCT	TCTTAGGGAA
7021	CAGCTGGGGA	CAGAGAACCA	CAGACACTCT	GAGGAGCCAC	CCTGAGGCCT	CTTTTGCCAG
7081	AGGACCTTAC	AGCCTCCCTG	GCAGCAGTTC	CGCCAGCATT	TCTGTAAATG	CCCTCATGCC
7141	AGGGTGCGCG	CCGGCTGTCA	GCACGAGAGG	GACGTTGGTC	TGTCCCCTGG	CACCGAGTCA
7201	GTCAGAAGGG	TGGCCAGGGC	CCCCTTGGGC	CCCTCCAGAG	ACAATCCACT	GTGGTCACAC
7261	GGCTCGGTGG	CAGGAAGTGC	TGTTCTCTGCA	GCTGTGGGGA	CAGGGAGTGT	GGATGAAGCC
7321	AGGCTGGGTT	TGTCTGAAGA	CGGAGGCCCC	GAAAGGTGGC	AGCCTGGCCT	ATAGCAGCAG
7381	CAACTCTTGG	ATTTATTTGA	AAGATTTTCT	TCACGGTTCT	GAGTCTTGGG	GGTGTTAGAG
7441	GCTCAGAACC	AGTCCAGCCA	GAGCTCTGTC	ATGGGCACGT	AGACCCGGTC	CCAGGGCCTT
7501	TGCTCTTTGC	TGTCCTCAGA	GGCCTCTGCA	AAGTAGAAAC	AGGCAGCCTT	GTGAGTCCCC
7561	TCCTGGGAGC	AACCAACCCCT	CCCTCTGAGA	TGCCCCGGGG	CCAGGTACAG	TGTGGTGAAA
7621	GGTAGGGATG	CAGCCAGCTC	AGGGAGTGGC	CCAGAGTTCC	TGCCCCACCA	AGGAGGCTCC
7681	CAGGAAGGTC	AAGGCACCTG	ACTCTGGGCG	TGCTTCCCTC	CCCTCCCCCT	CCCAGGTCAG
7741	GAAGGTGGGA	AAGGGCTGGG	GTGTCTGTGA	CCCTGGCAGT	CACTGAGAAG	CAGGGTGGAA
7801	GCAGCCCCCT	GCAGCACGCT	GGGTCACTGG	TCTTACCAGA	TGGATACGCA	GCAACTCTCT
7861	TTTGAACCTT	TTTATTTTCC	TGGCAGGAAG	AAGAGGGATC	CAGCAGTGAG	ATCAGGCAGG
7921	TTCTGTGTTG	CACAGACAGG	GAAACAGGCT	CTGTCCACAC	AAAGTCGGTG	GGGCCAGGAT
7981	GAGGCCAGT	CTGTTACAC	ATGGCTGCTG	CCTCTCAGCT	CTGCACAGAC	GTCTCTGCTC



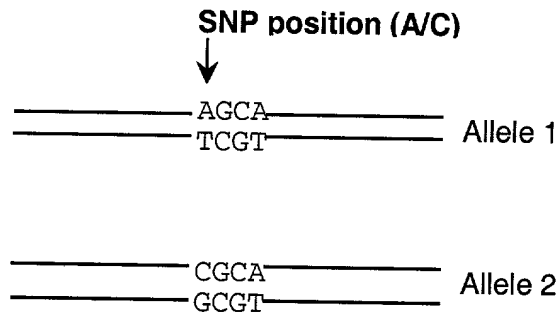
FIGURE 2

1 GAATTCAAGA CCAGCCTGGA CAACTTGGAA GAACCSGGTC TCTACAAAA ATACAAAATT  
61 AGCTGGGATT GGGTGCGGTG GCTCATGCCT ATAATCCCAG CACTTTGGGA GCCTGAGGTG  
121 GGTGGATCAC CTGAAGTCAG GAGTTCAAGA CTAGCCTGGC CAACATGGTG AAACCCTATC  
181 TCTACTGAAA ATAYAAAAAG CTAGACGTGG TGGCACACAC CTGTAATCCC AGCTACTTAG  
241 GAGGCTGAGG CAGGAGAATT GCTTGAAGCC TAGAGGTGAA GGTTGTAGTG AGCCGAGATT  
301 GCATCATTGC ACAATGGAGG GGAGCCACCA GCCTGGGCAA CAAGAGGAAA TCTCCGTCTC  
361 CAAAAAAAAA AAAAAAAAAA AAAGRATTAG GCTGGGTGGT GCCTGTAGTC CCAGCTACTT  
421 GGGAGGCAGG GGGTCCACTT GATGTCGAGA CTGCAGTGAG CCATGATCCT GCCACTGCAC  
481 TCCGGCCTGG GCAACAGAGT GAGACCCTGT CTAAAGAAAA AAAAAATAAA GCAACATATC  
541 CTGAACAAAG GATCCTCCAT AACGTTCCCA CCAGATTTCT AATCAGAAAC ATGGAGGCCA  
601 GAAAGCAGTG GAGGAGGACR ACCCTCAGGC AGCCCGGGAG GATGTTGTCA CAGGCTGGGG  
661 CAAGGGCCTT CCGGCTACCA ACTGGGAGCT CTGGGAACAG CCCTGTTGCA AACAAGAAGC  
721 CATAGCCCGG CCAGAGCCCA GGAATGTGGG CTGGGCTGGG AGCAGCCTCT GGACAGGAGT  
781 GGTCCCATCC AGGAAACCTC CGGCATGGCT GGGAAAGTGGG GTACTTGGTG CCGGGTCTGT  
841 ATGTGTGTGT GACTGGTGTG TGTGAGAGAG AATGTGTGCY CTAAGTGTC GTGTGAGTCT  
901 GTGTATGTGT GAATATTGTC TTTGTGTGGG TGATTTCTG CRTGTGTAAT CGTGTCCCTG  
961 CAAGTGTGAA CAAGTGGACA AGTGTCTGGG AGTGGACAAG AGATCTGTGC ACCATCAGGT  
1021 GTGTGCATAG CGTCTGTGCA TGTCAAGAGT GCAAGGTGAA GTGAAGGGAC CAGGCCCATG  
1081 ATGCCACTCA TCATCAGGAG CTCTAAGGCC CCAGGTAAGT GCCAGTGACA GATAAGGGTG  
1141 CTGAAGGTCA CTCTGGAGTG GGCAGGTGGG GGTAGGGAAA GGGCAAGGCC ATGTTCTGGA  
1201 GGAGGGGTG TGACTACATT AGGGTGTATG AGCCTAGCTG GGAGGTGGAT GGCCRGGTCC  
1261 ACTGAAACCC TGGTTATCCC AGAAGGCTTT GCAGGCTTCA GGAGCTTGA GTGGGGAGAG  
1321 GGGGTGACTT CTCCGACCAG GCCCTCCAC CGGCCTACCC TGGGTAAGGG CCTGGAGCAG  
1381 GAAGCAGGGG CAAGAACCTC TGGAGCAGCC CATACCCGCC CTGGCCTGAC TCTGCCACTG  
1441 GCAGCACAGT CAACACAGCA GGTTCACTCA CAGCAGAGGG CAAAGGCCAT CATCAGCTCC  
1501 CTTTATAAGG GAAGGGTCAC GCGCTCGGTG TGCTGAGAGT GTCCTGCCTG GTCCTCTGTG  
1561 CCTGGTGGGG TGGGGGTGCC AGGTGTGTCC AGAGGAGCCC ATTTGGTAGT GAGGCAGGTA  
1621 TGGGGCTAGA AGCACTGGTG CCCCTGGCCG TGATAGTGGC CATCTTCCTG CTCCTGGTGG

T06280"0T24550

FIGURE 3

## One Base Sequencing (OBS) Outline



Add Cy5-ddATP + dTTP,dCTP,dGTP + DNA polymerase

